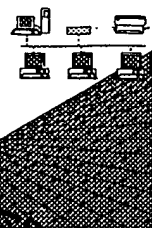


CRFI

BIOTECHNOLOGY
SYSTEMS
BRANCH



AUG 11 2003

TECH CENTER 1600/2900

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/001, 934 A
Source: 1600
Date Processed by STIC: 8-4-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/001,934A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ✓ Use of <220> Sequence(s) 21, 88, 94 *these may be others* missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



Does Not Comply
Corrected Diskette Needed

1600

RAW SEQUENCE LISTING

DATE: 08/04/2003

PATENT APPLICATION: US/10/001,934A

TIME: 07:19:03

Input Set : N:\efs\10001934A\GPCG-P01-003SequenceListing.txt

Output Set: N:\CRF4\08012003\J001934A.raw

4 <110> APPLICANT: NAGY et al.
6 <120> TITLE OF INVENTION: HUMAN POLYPEPTIDES CAUSING OR LEADING TO THE KILLING
7 OF CELLS INCLUDING LYMPHOID TUMOR CELLS
9 <130> FILE REFERENCE: GPCG-P01-003
11 <140> CURRENT APPLICATION NUMBER: 10/001934A
12 <141> CURRENT FILING DATE: 2001-11-15
14 <150> PRIOR APPLICATION NUMBER: PCT/US01/15625
15 <151> PRIOR FILING DATE: 2001-05-14
17 <160> NUMBER OF SEQ ID NOS: 94
19 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

318 <210> SEQ ID NO: 21
319 <211> LENGTH: 10
320 <212> TYPE: PRT
321 <213> ORGANISM: artificial sequence
323 <220> FEATURE:
W--> 324 <221> NAME/KEY: MS-GPC8-6-VH-CDR3
325 <222> LOCATION: (1)..(10)
W--> 327 <223> OTHER INFORMATION:
W--> 327 <400> 21
329 Ser Pro Arg Tyr Arg Gly Ala Phe Asp Tyr
E--> 330 1 10
2160 <210> SEQ ID NO: 88
2161 <211> LENGTH: 115
2162 <212> TYPE: PRT
2163 <213> ORGANISM: artificial sequence
2165 <220> FEATURE:
W--> 2166 <221> NAME/KEY: MS-GPC8-27-VL
2167 <222> LOCATION: (1)..(115)
W--> 2169 <223> OTHER INFORMATION:
W--> 2169 <400> 88
2170 Asp Ile Val Leu Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
2171 1 5 10 15
2173 Arg Val Thr Ile Ser Cys Ser Gly Ser Ser Asn Ile Gly Ser Asn
2174 20 25 30
2176 Tyr Val Ser Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
2177 35 40 45
2179 Ile Tyr Asp Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe
2180 50 55 60
2182 Ser Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Thr Gly Leu

See page 3

When using Artificial Sequence
for numeric identifier (213). The
Source of the genetic material must
be provided in (223). See item # 11 on
Error Summary Sheet.

Amino Acids should be
numbered every 5.

Same as above. Note there
may be other sequences with this
problem. See page 3.

No blank
spaces should be
included in
numbering count.

RAW SEQUENCE LISTING

DATE: 08/04/2003

PATENT APPLICATION: US/10/001,934A

TIME: 07:19:03

Input Set : N:\efs\10001934A\GPCG-P01-003SequenceListing.txt

Output Set: N:\CRF4\08012003\J001934A.raw

E--> 2183 (65) (70) (75) (80)
 2185 Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Met Asn
 E--> 2186 (85) (90) (95)
 2188 Val His Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
 E--> 2189 (100) (110) (115)
 2364 <210> SEQ ID NO: 94
 2365 <211> LENGTH: 109
 2366 <212> TYPE: PRT
 2367 <213> ORGANISM: artificial sequence
 2369 <220> FEATURE:
 W--> 2370 <221> NAME/KEY: MS-GPC8-27-41-VL
 2371 <222> LOCATION: (1)..(108)
 W--> 2373 <223> OTHER INFORMATION:
 W--> 2373 <400> 94
 2374 Asp Ile Val Leu Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
 2375 1 5 10 15
 2377 Arg Val Thr Ile Ser Cys Ser Gly Ser Glu Ser Asn Ile Gly Asn Asn
 2378 20 25 30
 2380 Tyr Val Gln Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
 2381 35 40 45
 2383 Ile Tyr Asp Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
 2384 50 55 60
 2386 Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Thr Gly Leu Gln
 2387 65 70 75 80
 2389 Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Met Asn Val
 2390 85 90 95
 2392 His Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
 2393 100 105
 E--> 2397 1

Numbering
must be every 5.

Same as first page
see page 3

Remove extra material at end of
file.

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/001,934A

DATE: 08/04/2003
TIME: 07:19:04

Input Set : N:\efs\10001934A\GPCG-P01-003SequenceListing.txt
Output Set: N:\CRF4\08012003\J001934A.raw

Use of <220> Feature (NEW RULES):

Sequence(s) __ are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29

Seq#:30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53

Seq#:54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75,76,77

Seq#:78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94

VERIFICATION SUMMARY

DATE: 08/04/2003

PATENT APPLICATION: US/10/001,934A

TIME: 07:19:04

Input Set : N:\efs\10001934A\GPCG-P01-003SequenceListing.txt

Output Set: N:\CRF4\08012003\J001934A.raw

L:27 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:34 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:43 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:60 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:63 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:3, <213>
ORGANISM:artificial sequence
L:63 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:63
L:74 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:77 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:4, <213>
ORGANISM:artificial sequence
L:77 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:77
L:87 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:90 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:5, <213>
ORGANISM:artificial sequence
L:90 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:90
L:100 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:103 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:6, <213>
ORGANISM:artificial sequence
L:103 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:6,Line#:103
L:115 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7
L:118 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:7, <213>
ORGANISM:artificial sequence
L:118 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:7,Line#:118
L:129 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:132 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:8, <213>
ORGANISM:artificial sequence
L:132 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:8,Line#:132
L:144 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9
L:147 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:9, <213>
ORGANISM:artificial sequence
L:147 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:9,Line#:147
L:159 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:162 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:10, <213>
ORGANISM:artificial sequence
L:162 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10,Line#:162
L:174 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11
L:177 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:11, <213>
ORGANISM:artificial sequence
L:177 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:177
L:189 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:12
L:192 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:12, <213>
ORGANISM:artificial sequence
L:192 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:12,Line#:192
L:204 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:13
L:207 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:13, <213>
ORGANISM:artificial sequence
L:207 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:207
L:218 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:221 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:artificial sequence
L:221 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:221
L:233 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15

L:236 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
ORGANISM:artificial sequence
L:236 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:236
L:249 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:252 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:16, <213>
ORGANISM:artificial sequence
L:252 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:252
L:264 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:267 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>
ORGANISM:artificial sequence

VERIFICATION SUMMARY

DATE: 08/04/2003

PATENT APPLICATION: US/10/001,934A

TIME: 07:19:04

Input Set : N:\efs\10001934A\GPCG-P01-003SequenceListing.txt

Output Set: N:\CRF4\08012003\J001934A.raw

L:267 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:17,Line#:267
L:280 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:18
L:283 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:18, <213>
ORGANISM:artificial sequence
L:283 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:18,Line#:283
L:294 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:19
L:297 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:19, <213>
ORGANISM:artificial sequence
L:297 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:19,Line#:297
L:308 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:20
L:311 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:20, <213>
ORGANISM:artificial sequence
L:311 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:20,Line#:311
L:324 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:21
L:327 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:21, <213>
ORGANISM:artificial sequence
L:327 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:21,Line#:327
L:330 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:21
L:339 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:22
L:342 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:22, <213>
ORGANISM:artificial sequence
L:342 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:22,Line#:342
L:354 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:23
L:357 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:23, <213>
ORGANISM:artificial sequence
L:357 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:23,Line#:357
L:369 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:24
L:372 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:24, <213>
ORGANISM:artificial sequence
L:372 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:24,Line#:372
L:383 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:25
L:386 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:25, <213>
ORGANISM:artificial sequence
L:386 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:25,Line#:386
L:397 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:26
L:400 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:26, <213>
ORGANISM:artificial sequence
L:400 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:26,Line#:400
L:412 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:27
L:415 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:27, <213>
ORGANISM:artificial sequence
L:415 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:27,Line#:415
L:427 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:28
L:442 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:29
L:457 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:30
L:471 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31
L:485 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:32
L:501 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:33
L:516 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:34
L:531 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:35
L:545 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:36
L:559 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:37
L:574 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:38

L:588 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:39
L:603 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:40
L:617 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:41
L:633 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:42
L:648 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:43
L:663 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:44

VERIFICATION SUMMARY

DATE: 08/04/2003

PATENT APPLICATION: US/10/001,934A

TIME: 07:19:04

Input Set : N:\efs\10001934A\GPCG-P01-003SequenceListing.txt

Output Set: N:\CRF4\08012003\J001934A.raw

L:678 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:45
L:692 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:46
L:707 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:47
L:721 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:48
L:736 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:49
L:751 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:50
L:2183 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:88
M:332 Repeated in SeqNo=88
L:2189 M:252 E: No. of Seq. differs, <211> LENGTH:Input:115 Found:109 SEQ:88
L:2397 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:94